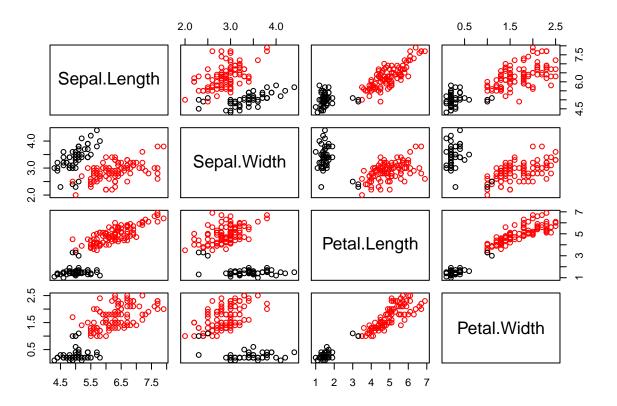
homework6.R.

```
library(plyr)
library(e1071)
library(caret)
## Loading required package: lattice
## Loading required package: ggplot2
library(fpc)
data <- read.table('iris.data',sep=",",header=F)</pre>
names (data)
## [1] "V1" "V2" "V3" "V4" "V5"
set.seed(666)
names(data) <-c('sepalLength','sepalWidth','petalLength','petalWidth','category')</pre>
CVgroup <- function(k,datasize,seed){</pre>
  cvlist <- list()</pre>
  set.seed(seed)
  n <- rep(1:k,ceiling(datasize/k))[1:datasize] #divide data into k, qenerate whole dataset n
  temp <- sample(n,datasize)</pre>
                                #shuffle n
  x \leftarrow 1:k
  dataseq <- 1:datasize</pre>
  cvlist <- lapply(x,function(x) dataseq[temp==x]) #randomly generate k random sequential dataset</pre>
  return(cvlist)
}
k <- 10
datasize <- nrow(iris)</pre>
cvlist <- CVgroup(k=k, datasize = datasize, seed=1206)</pre>
train <- iris[-cvlist[[1]],]</pre>
test <- iris[cvlist[[1]],]</pre>
nb1 <- naiveBayes(Species ~., data=train)</pre>
prediction <- predict(nb1, test)</pre>
confusionMatrix <- table(prediction,test$Species)</pre>
confusionMatrix
##
## prediction setosa versicolor virginica
```

```
0
##
     setosa
                    4
                               0
                     0
##
    versicolor
                                3
                                          0
                                          7
     virginica
                     0
accuracy = sum(confusionMatrix[row(confusionMatrix)==col(confusionMatrix)]) / sum(confusionMatrix)
accuracy
## [1] 0.9333333
class = sort(unique(prediction))
tp=NA
fp=NA
fn=NA
tn=NA
for(i in 1:length(class)){
  tp[i] = sum(test$Species==class[i] & prediction==class[i])
  tn[i] = sum(test$Species!=class[i] & prediction!=class[i])
  fp[i] = sum(test$Species==class[i] & prediction!=class[i])
  fn[i] = sum(test$Species!=class[i] & prediction==class[i])
precision = tp/(tp + fp)
recall = tp/(tp + fn)
f1score = (2*precision*recall)/(precision*recall)
names(f1score) = class
names(precision) = class
names(recall) = class
print(table(test$Species,prediction))
##
               prediction
##
                setosa versicolor virginica
##
                   4
                               0
                                          0
    setosa
                                          1
                     0
                                3
##
     versicolor
                               0
##
     virginica
                     0
print('precision')
## [1] "precision"
print(precision)
##
       setosa versicolor virginica
##
         1.00
                  0.75
                               1.00
print('recall')
## [1] "recall"
```

```
print(recall)
##
       setosa versicolor virginica
##
       1.000
                  1.000
                               0.875
print('f1score')
## [1] "f1score"
print(f1score)
       setosa versicolor virginica
## 1.0000000 0.8571429 0.9333333
print('mean(f1score)')
## [1] "mean(f1score)"
print(mean(f1score))
## [1] 0.9301587
#Qusetion 4
newIris<-iris[1:4]</pre>
kc<-kmeans(newIris,2)</pre>
kc$tot.withinss
## [1] 152.348
kc$betweenss
## [1] 529.0226
plot(newIris, col=kc$cluster)
```

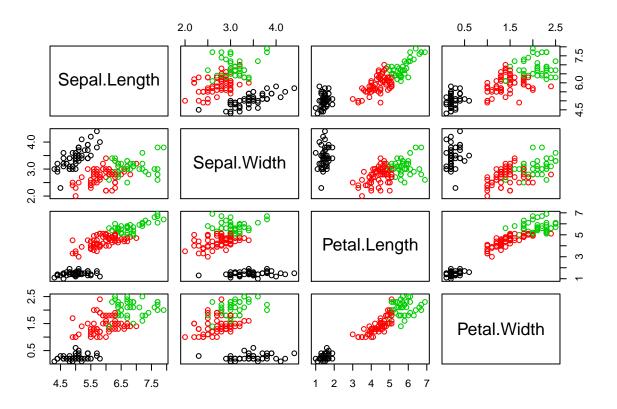


kc<-kmeans(newIris,3)
kc\$tot.withinss</pre>

[1] 78.85144

kc\$betweenss

[1] 602.5192

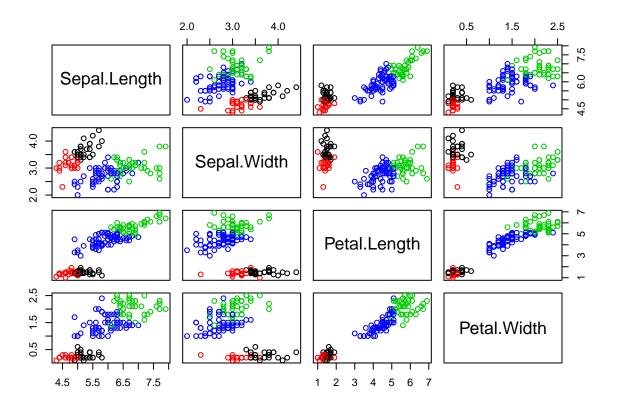


kc<-kmeans(newIris,4)
kc\$tot.withinss</pre>

[1] 71.44525

kc\$betweenss

[1] 609.9254

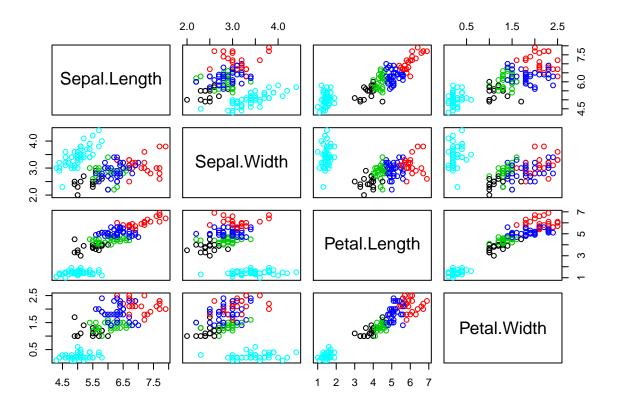


kc<-kmeans(newIris,5)
kc\$tot.withinss</pre>

[1] 51.08942

kc\$betweenss

[1] 630.2812

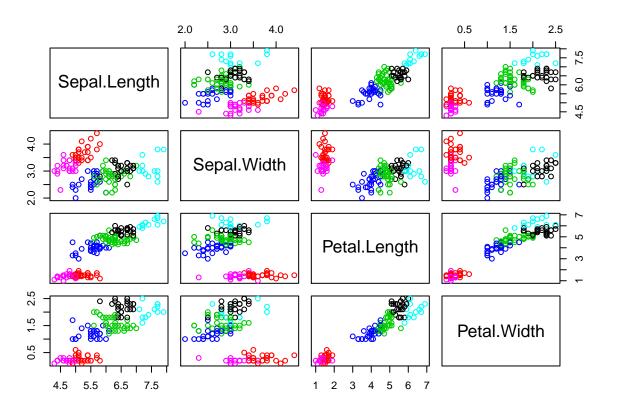


kc<-kmeans(newIris,6)
kc\$tot.withinss</pre>

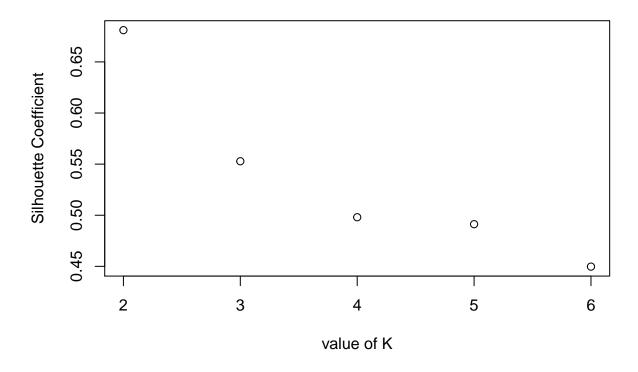
[1] 39.05498

kc\$betweenss

[1] 642.3156



```
K <- 2:6
round <- 1
rst <- sapply(K,function(i){
   mean(sapply(1:round,function(r){
      result <- kmeans(newIris, i)
      stats <- cluster.stats(dist(newIris), result$cluster)
      stats$avg.silwidth
   }))
})
plot(K,rst,type='p', xlab='value of K',ylab='Silhouette Coefficient')</pre>
```



```
#rbind(K,rst)
kc <- kmeans(newIris,3)</pre>
confusion_matrix <- table(kc$cluster,iris$Species)</pre>
confusion_matrix
##
       setosa versicolor virginica
##
##
     1
            17
##
     2
            33
                         0
                                    0
     3
             0
                        46
                                   50
##
```